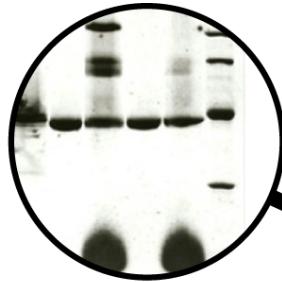
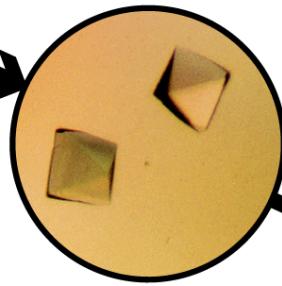


High-throughput Crystallography

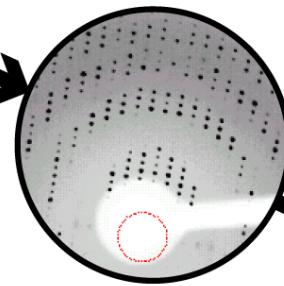


Automated Protein Purification



Automated Crystallization

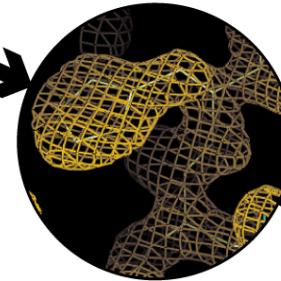
- The structural database must expand
- NIH funding centers to collectively solve several 1000 structures in next 5 years



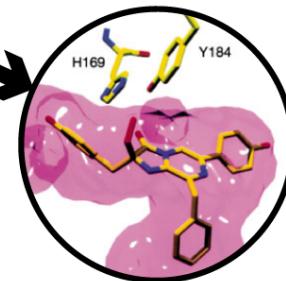
Automated Data Collection

PHENIX

*Python-based
Hierarchical
ENvironment for
Integrated
Xtallography*



Automated Structure Solution



Interpretation

Structure solution will take hours - days

Technology benefits all crystallographers

<http://www.phenix-online.org>



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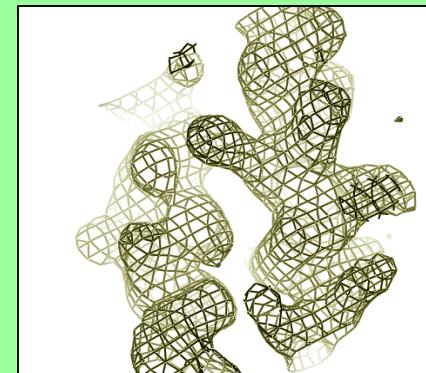


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The *PHENIX* project

- Goal: develop new crystallographic software for automated structure determination, using modern algorithms and advances in computer science
- The new system will be comprehensive and flexible enough to automate structure solution, but allow user interaction, input and control
- Automated crystallography at medium to low-resolution (3Å)
 - Maximum Likelihood methods
 - Phasing
 - Molecular replacement
 - Density modification
 - Refinement
 - Pattern matching for automated building
 - Simulated annealing refinement



The *PHENIX* project

A collaboration between several groups

Computational Crystallography Initiative (LBNL)

- Paul Adams, Ralf Grosse-Kunstleve*
- Nigel Moriarty, Nicholas Sauter, DK Smith*



Los Alamos National Lab (LANL)

- Tom Terwilliger, Li-Wei Hung*



Cambridge University

- Randy Read, Airlie McCoy, Laurent Storoni*



Texas A&M University

- Tom Ioerger, Jim Sacchettini, Kreshna Gopal, Tod Romo*



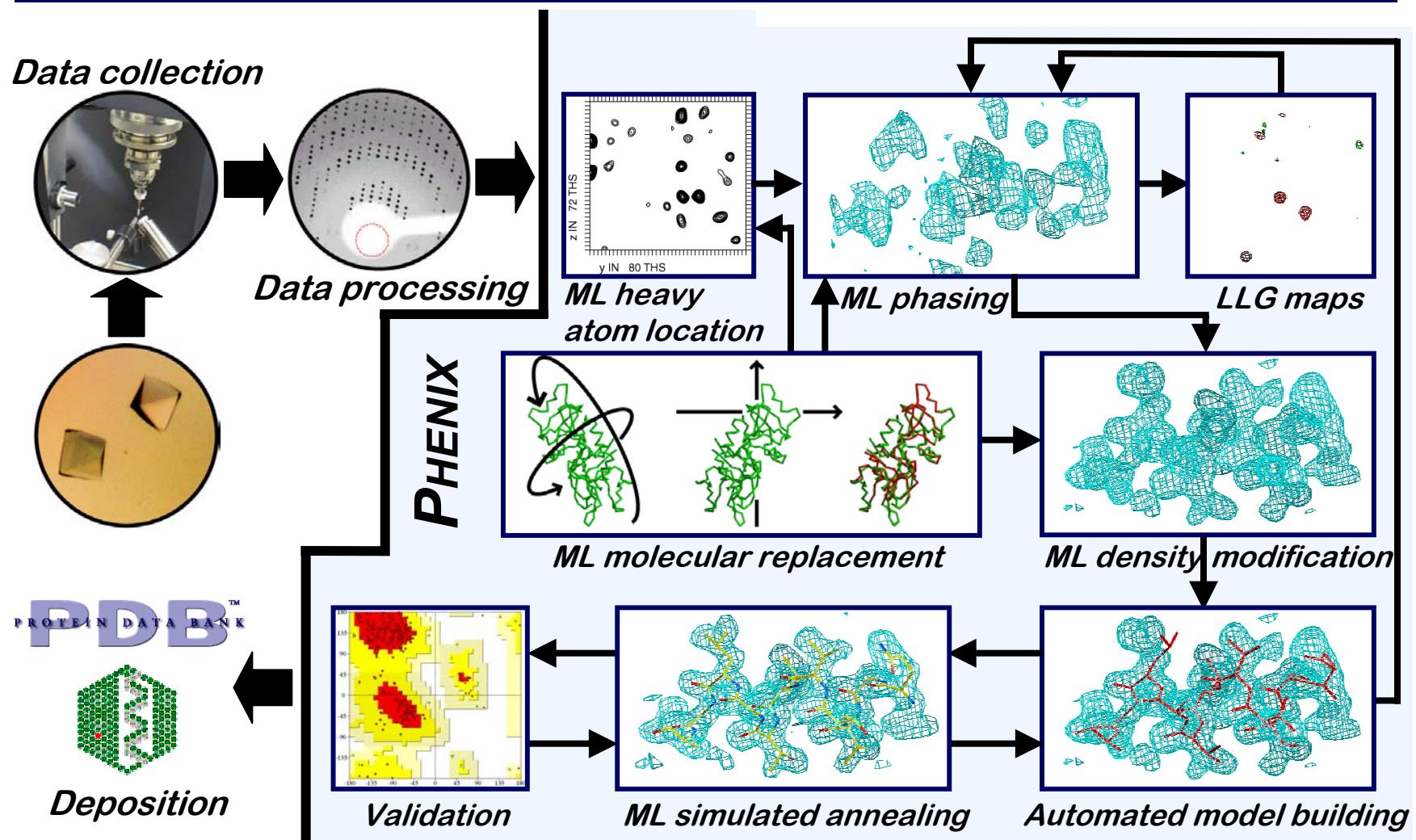
Funding: NIH (NIGMS, PSI) Program Project (LBNL lead, PDA director)



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Automated Structure Solution



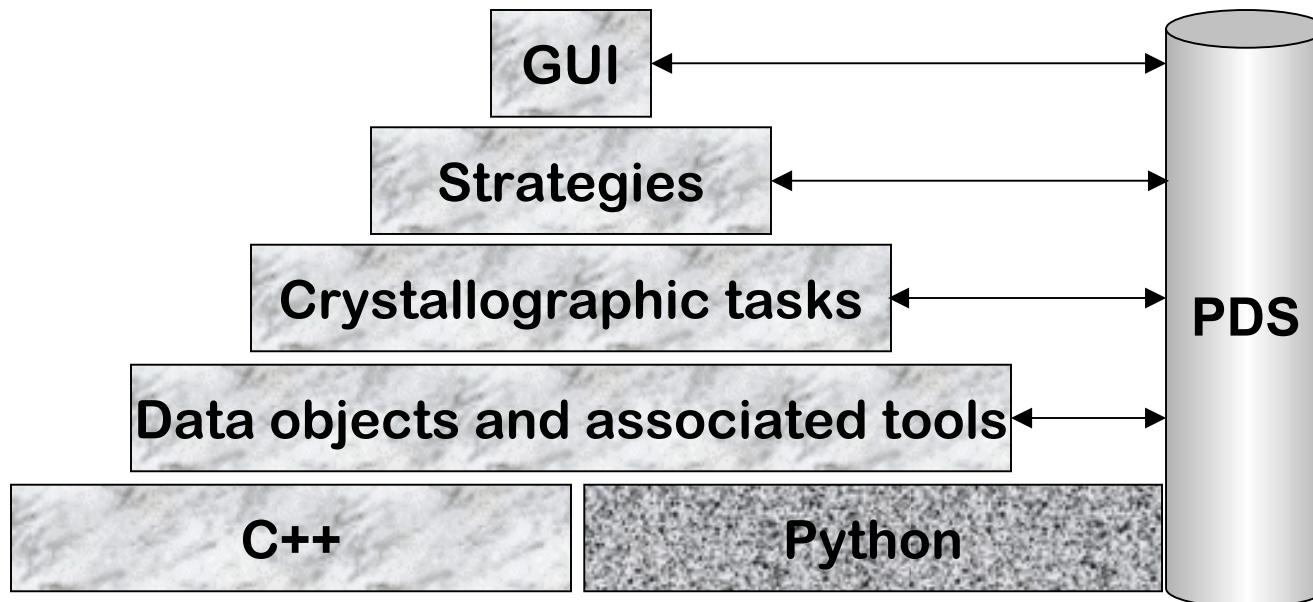
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ATM

A Collaborative Framework

- Flexible: can automate the whole structure solution process but also allow user input and control
- Built around a scripting language: efficient development of new ideas



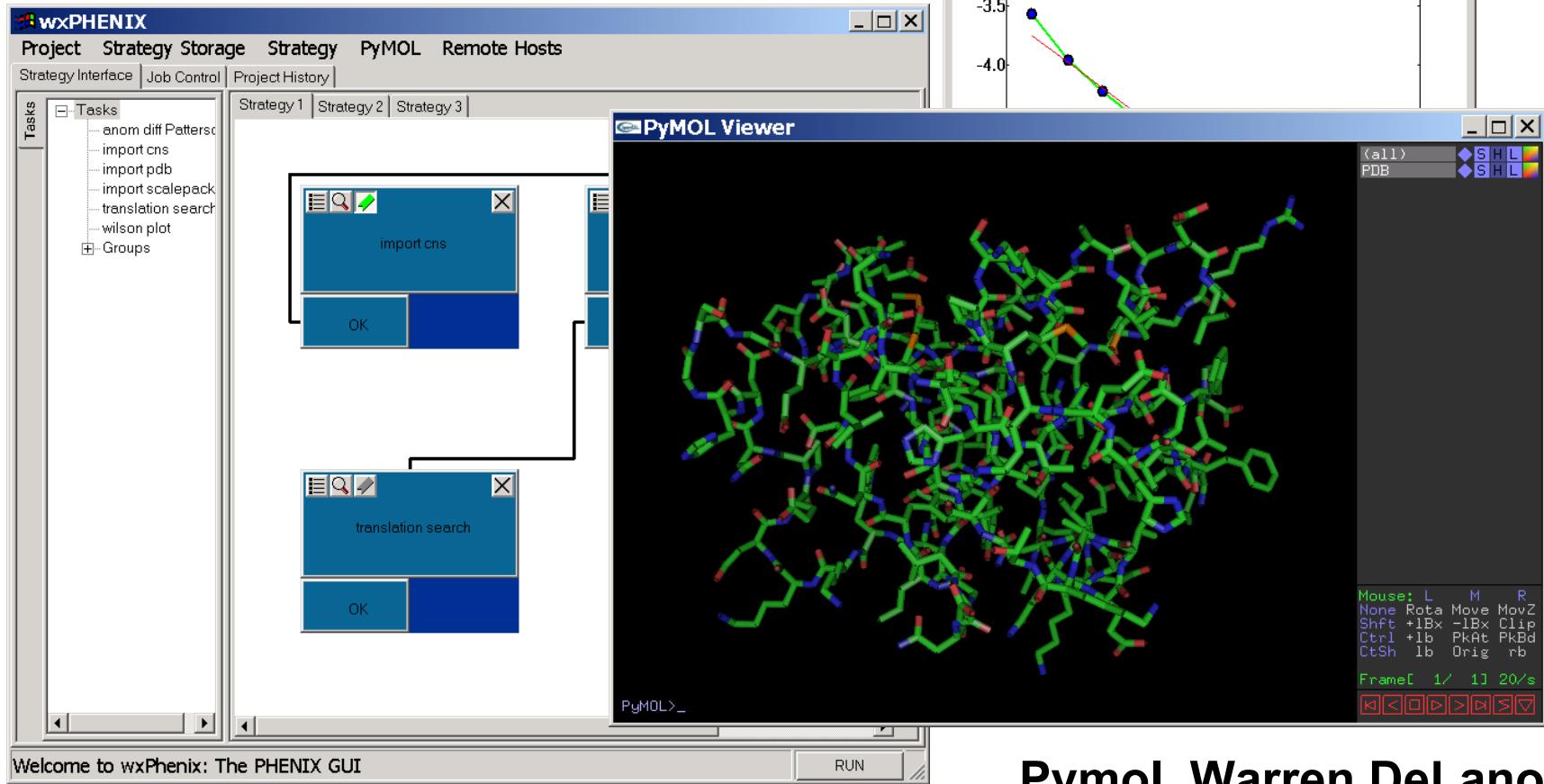
- Project Data Storage (PDS): for each structure maintains a history of the structure solution and all the data generated.

PHENIX Infrastructure

- **Basic objects:**
 - Structure factors
 - Map
 - Coordinates
- **Visual programming**
 - Strategies
- **Project Data Storage**
 - Type based data storage
- **Project history browser**
 - Linked to visualization tools
- **Distributed computing**
- **Modularity**
 - Object oriented programming
 - Dynamically linked objects
- **Scripting language**
 - Python
- **Open Source cctbx project**
 - Available to all developers
- **Source code distribution to academic groups**
- **Other developers can build “plug-ins” that make use of *PHENIX***

Interfacing with the GUI

- Can use other programs as helper applications to view results



Welcome to wxPhenix: The PHENIX GUI

RUN

Pymol, Warren DeLano



PHENIX



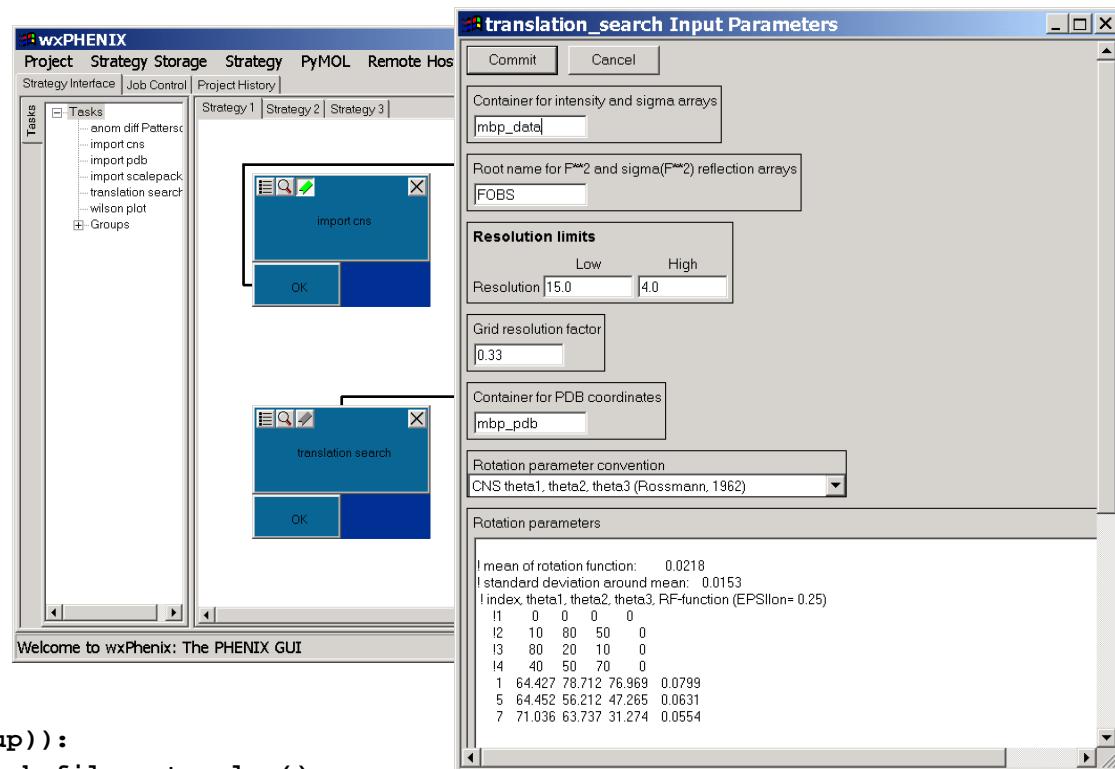
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Developing new Strategies and Tasks

- Visual programming allows easy creation of complex algorithms
- GUI elements are constructed from Python objects (created in text files)
- Python task files could call other programs

```
def __init__(self):  
    ...  
  
    task_parameters = parameters()  
    group(  
        ...  
        name = 'file_group',  
        ...  
        parameters = parameters()  
        define(  
            ...  
            name = 'scalepack_file',  
            ...  
        ),  
        ),  
    )  
  
    task_branches = {...}  
  
def Run(self):  
    inp = self.get_parameters()  
  
    for i in xrange(len(inp.file_group)):  
        print inp.file_group[i].scalepack_file.get_value()
```



Implementing algorithms - Python

#READ INTENSITY DATA

```
S = xarray.ScalepackReader(scalepackfile)
A = S.newlspace()
S.loadlspace(A,"lobs")
```

#CALCULATE AMPLITUDES

```
lobs = A.use_IntensityArray("lobs")
F = lobs.F()
sigF = lobs.sigF()
```

#SELECTION ON SIGMA (F)

```
S1 = xarray.greater(F, SigCut*sigF)
```

#CALCULATE ANOMALOUS DIFFERENCES

```
anom_diff = F.plus() - F.minus()
sigdiff = xarray.sqrt(xarray.pow2(sigF.plus())+xarray.pow2(sigF.minus()))
```

#SELECTION ON SIGMA (delta F)

```
S3 = xarray.greater(xarray.abs(anom_diff), DifCut * sigdiff)
```

#CALCULATE E VALUES

```
ebin = A.Binner(8, 4.0)
amp = xarray.sqrt(dfsquare)
E = amp / ebin.average(amp)
Esquare = E * E
```

#REMOVE ORIGIN

```
Esq_rem = Esquare - ebin.saverage(Esquare)
```

#CALCULATE AND OUTPUT PATTERSON MAP

```
proform = xarray.MapForm(A,0,4.0,0.25) #0=HERMITIAN, 1=NONHERMITIAN
pro = xarray.Xarray(proform,1)          #0=RealSpace, 1=ReciprocalSpace
pro.loadArray(Esq_rem)
pro.toRealSpace()
pro.real2CNSfile(outputfile)
```

Python Script

- The *PHENIX* scripting language uses Python, extended with C++
- Provides the convenience of a scripting language with the performance of a compiled language
- Many ideas can be tested first using Python, then optimized as needed
- The *PHENIX* scripting language provides access to crystallographic data objects, and tools (e.g. structure factors and structure factor algebra)
- No compilation is required to run this script



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Implementing algorithms – C++

//READ INTENSITY DATA

```
ScalepackReader S = ScalepackReader(scalepackfile);
Ispace A = S.newIspace();
S.loadIspace(A,"lobs");
```

//CALCULATE AMPLITUDES

```
IntensityArray lobs = A.use<IntensityArray>("lobs");
IspaceArray F = lobs.F();
IspaceArray sigF = lobs.sigF();
```

//SELECTION ON SIGMA (F)

```
IspaceArray S1 = Xalgebra::greater(F, SigCut*sigF);
```

//CALCULATE ANOMALOUS DIFFERENCES

```
IspaceArray anom_diff = F.plus() - F.minus();
IspaceArray sigdiff = sqrt (pow2(sigF.plus()) + pow2(sigF.minus()));
```

//SELECTION ON SIGMA (delta F)

```
IspaceArray S3 = Xalgebra::greater(abs(anom_diff), DifCut * sigdiff);
```

//CALCULATE E VALUES

```
Binner ebin = A.temporary<Binner>(8, 4.0);
IspaceArray amp = sqrt(dfsquare);
IspaceArray E = amp / ebin.average(amp);
IspaceArray Esquare = E * E;
```

//REMOVE ORIGIN

```
IspaceArray Esq_rem = Esquare - ebin.saverage(Esquare);
```

//CALCULATE AND OUTPUT PATTERSON MAP

```
MapForm proform(A, HERMITIAN, 4.0, 0.25);
Xarray pro(proform, ReciprocalSpace);
pro.loadArray(Esq_rem);
pro.toRealSpace();
pro.real2CNSfile(outputfile);
```

- C++ libraries implement crystallographic data objects and tools (xtbx, cctbx)
- C++ code can be used outside of the *PHENIX* framework
- The Boost.Python Library can be used to make the classes written in C++ available in Python
- Very similar coding with both C++ and Python

C++ Program



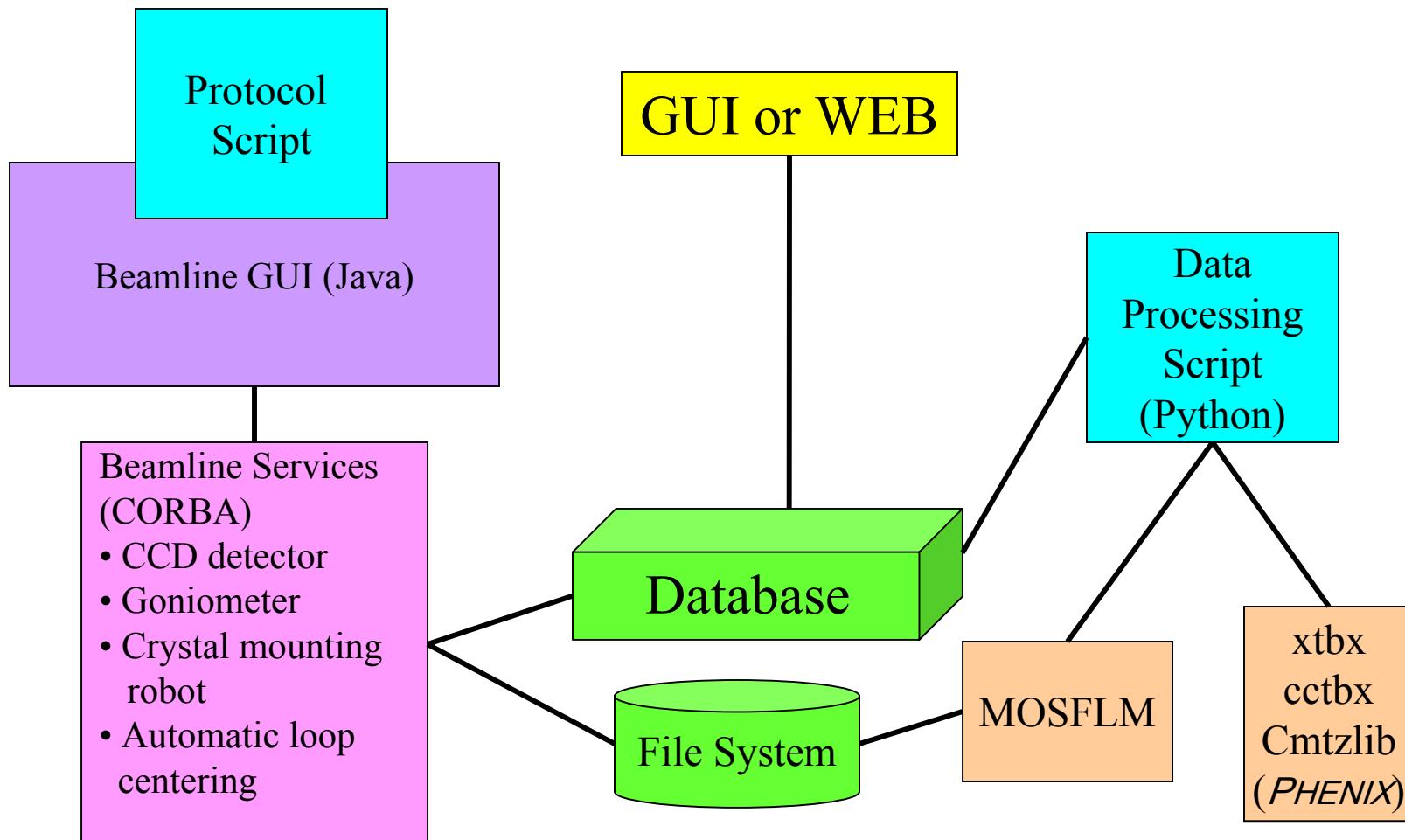
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Using *PHENIX* as an application



Nick Sauter and Thomas Earnest's group at the ALS

Summary

- The scope of the *PHENIX* software development is very broad:
 - New algorithms for structure solution
 - High level, generic tools (Strategy interface)
 - Crystallographic libraries (cctbx at SourceForge)
 - Low level “computer science” tools (Python/C++ interface)
- There are many opportunities for other developers to make use of the *PHENIX* infrastructure



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